

## NOTICE OF REFERENCES CITED

APPLICANT(S) : Ebner et al.

## U.S. PATENT DOCUMENTS

*	DOCUMENT NUMBER	DATE	NAME(S)	CLASS	SUBCLASS	FILING DATE
	5,194,596	3/16/93	Tischer et al.	530	399	
	5,256,766	10/26/93	Coughlin	-	-	
	58,242,798	9/7/93	Sutcliffe	-	-	

## FOREIGN PATENT DOCUMENTS

*	DOCUMENT NO.	DATE	COUNTRY	NAME	CLASS	SUBCLASS	PERTINENT DRW	SPEC
	01548	1/20/94	WO	Sibson	---	---		

## \* OTHER REFERENCES (INCLUDING AUTHOR, TITLE, DATE, PERTINENT PAGES, ETC.)

J. Skolnick et al., "From genes to protein structure and function: novel applications of computational approaches in the genomic era", Trends in Biotech 18(1):34-39, 2000.
P. Bork, "Powers and pitfalls in sequence analysis: The 70% hurdle", Genome Research 10:398-400, 2000.
T. Doerks et al., "Protein annotation: detective work for function prediction", Trends in Genetics, 4(g):248-250, June 1998.
T.F. Smith et al., "The challenges of genome sequence annotation or "The devil is in the details", Nature Biotechnology 15:1222-1223, 1997.
L. E. Benjamin et al., "A plasticity window for blood vessel remodelling is defined by pericyte coverage of the preformed endothelial network and is regulated by PDGF-B and VEGF." Development 125:1591-1598, 1998.
M. Bonaldo et al., Locus HUMOTIA, Genbank Accession number L23206 , accessed 3/1/01, Available 12/13/93.
L. Dumoutier et al., locus HSA277248, NCBI nucleic acid database, accessed 11/28/01, available 5/50/01.
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EXAMINER

DATE

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